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## Amendments to the Claims:

- 1. (Cancelled)
- 2. (Currently Amended) An isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:
  - a) a nucleotide sequence set forth in SEQ ID NO:1;
- b)—a nucleotide sequence that is an antisense sequence for the nucleotide sequence set forth in SEQ ID NO:1, wherein said antisense sequence hybridizes to the nucleotide sequence set forth in SEQ ID NO:1 under high stringency hybridization conditions of 50% formamide, 1 M NaCl, 1% SDS at 37°C, and a wash in 0.1X SSC at 60 to 65°C;
- e)b) a nucleotide sequence having at least-80% 95% sequence identity to the sequence set forth in SEQ ID NO:1, wherein said nucleotide sequence encodes a polypeptide having invertase inhibitor activity; and
- d)c) a nucleotide sequence that hybridizes to the nucleotide sequence set forth in SEQ ID NO:1 or a complement thereof under high stringency hybridization conditions of 50% formamide, 1 M NaCl, 1% SDS at 37°C, and a wash in 0.1X SSC at 60 to 65°C, or a complement thereof; and is a complement of a) or b).
- e)—a fragment of at least 50 contiguous nucleotides of the nucleotide sequence set forth in SEQ ID NO:1.
- 3. (Currently Amended) The nucleic acid molecule of claim 2, wherein said sequence encodes an amino acid selected from the group consisting of:
  - the amino acid sequence set forth in SEQ ID NO: 2[[.]]; and
- b) an amino sequence having 95% sequence identity to the sequence set forth in SEQ ID NO:2.
- 4. (Previously Presented) A chimeric gene comprising a plant-functional promoter operably linked to the nucleotide sequence of claim 2.

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- 5. (Previously Presented) The chimeric gene of claim 4, wherein the nucleotide sequence encodes the amino acid sequence set forth in SEQ ID NO:2.
- 6. (Previously Presented) The chimeric gene of claim 4, wherein said nucleotide sequence is the sequence set forth in SEQ ID NO:1.
- 7. (Currently Amended) The chimeric gene of claim 4, wherein said nucleotide sequence is the an antisense sequence of the sequence set forth in SEQ ID NO:1, wherein said antisense sequence hybridizes to the nucleotide sequence set forth in SEQ ID NO:1 under high stringency hybridization conditions of 50% formamide, 1 M NaCl, 1% SDS at 37°C, and a wash in 0.1X SSC at 60 to 65°C.
  - 8. (Original) A vector comprising the chimeric gene of claim 4.
  - 9. (Original) A plant cell transformed with the chimeric gene of claim 4.
  - 10. (Original) A plant comprising the chimeric gene of claim 4.
- 11. (Currently Amended) A transformed plant having incorporated into its genome a DNA molecule, said molecule comprising a nucleotide sequence operably linked to a promoter capable of driving expression of a gene in a plant cell, wherein said nucleotide sequence is selected from the group consisting of:
- a) a sequence encoding an invertase inhibitor having the amino acid sequence set forth in SEQ ID NO:2;
  - b) the nucleotide sequence set forth in SEQ ID NO:1;
- e) a nucleotide sequence that is an antisense sequence for the nucleotide sequence set forth in SEQ ID NO:1, wherein said antisense sequence hybridizes to the nucleotide

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sequence set forth in SEQ ID NO:1 under high stringency hybridization conditions of 50% formamide, 1 M NaCl, 1% SDS at 37°C, and a wash in 0.1X SSC at 60 to 65°C;

- d)c) a nucleotide sequence having at least-80% 95% sequence identity to the sequence set forth in SEQ ID NO:1, wherein said nucleotide sequence encodes a polypeptide having invertase inhibitor activity;
- e)d) a nucleotide sequence that hybridizes to the nucleotide sequence set forth in SEQ ID NO:1 or a complement thereof under high stringency hybridization conditions of 50% formamide, 1 M NaCl, 1% SDS at 37°C, and a wash in 0.1X SSC at 60 to 65°C, or a complement thereof; and encodes an amino acid sequence having 95% sequence identity to the sequence of SEQ ID NO:2; and
- f)e) a fragment of at least 50 contiguous nucleotides of the nucleotide sequence set forth in SEQ ID NO:1 nucleotide sequence that is a complement of any one of a) d).
- 12. (Previously Presented) The transformed plant of claim 11, wherein the nucleotide sequence encodes the amino acid sequence set forth in SEQ ID NO:2.
- 13. (Previously Presented) The transformed plant of claim 11, wherein the nucleotide sequence is the nucleotide sequence set forth in SEQ ID NO:1.
  - 14. (Cancelled)
  - 15. (Cancelled)
  - 16. (Original) The transformed plant of claim 11, wherein said plant is a dicot.
  - 17. (Original) The transformed plant of claim 11, wherein said plant is a monocot.
  - 18. (Original) The transformed plant of claim 17, wherein said plant is maize.

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- 19. (Original) Transformed seed of the plant of any one of claims 16-18.
- 20. (Currently Amended) A method for modulating invertase activity in a plant, said method comprising transforming said plant with a DNA construct, said construct comprising a promoter that drives expression in a plant cell operably linked with a nucleotide sequence selected from the group consisting of:
- a) a sequence encoding an invertase inhibitor having the amino acid sequence set forth in SEQ ID NO:2;
  - b) the nucleotide sequence set forth in SEQ ID NO:1;
- e) a nucleotide sequence that is an antisense sequence for the nucleotide sequence set forth in SEQ ID NO:1, wherein said antisense sequence hybridizes to the nucleotide sequence set forth in SEQ ID NO:1 under high stringency hybridization conditions of 50% formamide, 1 M NaCl, 1% SDS at 37°C, and a wash in 0.1X SSC at 60 to 65°C;
- d)c) a nucleotide sequence having at least-80% 95% sequence identity to the sequence set forth in SEQ ID NO:1, wherein said nucleotide sequence encodes a polypeptide having invertase inhibitor activity;
- e)d) a nucleotide sequence that hybridizes to the nucleotide sequence set forth in SEQ ID NO:1 or a complement thereof under high stringency hybridization conditions of 50% formamide, 1 M NaCl, 1% SDS at 37°C, and a wash in 0.1X SSC at 60 to 65°C, or a complement thereof; and encodes an amino acid sequence having 95% sequence identity to the sequence of SEQ ID NO:2; and
- f)e) a fragment of at least 50 contiguous nucleotides of the nucleotide sequence set forth in SEQ ID NO:1 nucleotide sequence that is a complement of any one of a) d).
- 21. (Currently Amended) A method for increasing seed yield in a plant, said method comprising transforming said plant with a DNA construct, said construct comprising a promoter

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that drives expression in a plant cell operably linked with a nucleotide sequence selected from the group consisting of:

- a) a sequence encoding-an invertase inhibitor having the amino acid sequence set forth in SEQ ID NO:2;
  - b) the nucleotide sequence set forth in SEQ ID NO:1;
- e) a nucleotide sequence that is an antisense sequence for the nucleotide sequence set forth in SEQ ID NO:1, wherein said antisense sequence hybridizes to the nucleotide sequence set forth in SEQ ID NO:1 under high stringency hybridization conditions of 50% formamide, 1 M NaCl, 1% SDS at 37°C, and a wash in 0.1X SSC at 60 to 65°C;
- d)c) a nucleotide sequence having at least-80% 95% sequence identity to the sequence set forth in SEQ ID NO:1, wherein said nucleotide sequence encodes a polypeptide having invertase inhibitor activity;
- e)d) a nucleotide sequence that hybridizes to the nucleotide sequence set forth in SEQ ID NO:1 or a complement thereof under high stringency hybridization conditions of 50% formamide, 1 M NaCl, 1% SDS at 37°C, and a wash in 0.1X SSC at 60 to 65°C, or a complement thereof; and encodes an amino acid sequence having 95% sequence identity to the sequence of SEQ ID NO:2; and
- f)e) a fragment of at least 50 contiguous nucleotides of the nucleotide sequence set forth in SEQ ID NO:1 nucleotide sequence that is a complement of any one of a) d).
- 22. (Currently Amended) A transformed plant cell having incorporated into its genome a DNA molecule, said molecule comprising a promoter capable of driving expression of a gene in a plant cell operably linked to a nucleotide sequence selected from the group consisting of:
- a) a sequence encoding an invertase inhibitor having the amino acid sequence set forth in SEQ ID NO:2;
  - b) the nucleotide sequence set forth in SEQ ID NO:1;

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e) a nucleotide sequence that is an antisense sequence for the nucleotide sequence set forth in SEQ ID NO:1, wherein said antisense sequence hybridizes to the nucleotide sequence set forth in SEQ ID NO:1 under high stringency hybridization conditions of 50% formamide, 1 M NaCl, 1% SDS at 37°C, and a wash in 0.1X SSC at 60 to 65°C;

- d)c) a nucleotide sequence having at least 80% 95% sequence identity to the sequence set forth in SEQ ID NO:1, wherein said nucleotide sequence encodes a polypeptide having invertase inhibitor activity;
- e)d) a nucleotide sequence that hybridizes to the nucleotide sequence set forth in SEQ ID NO:1 or a complement thereof under high stringency hybridization conditions of 50% formamide, 1 M NaCl, 1% SDS at 37°C, and a wash in 0.1X SSC at 60 to 65°C, or a complement thereof; and encodes an amino acid sequence having 95% sequence identity to the sequence of SEQ ID NO:2; and
- f)e) a fragment of at least 50 contiguous nucleotides of the nucleotide sequence set forth in SEQ ID NO:1 nucleotide sequence that is a complement of any one of a) d).
  - 23. (Cancelled)
  - 24. (Cancelled)
  - 25. (Cancelled)
  - 26. (Cancelled)